

09755071-010801

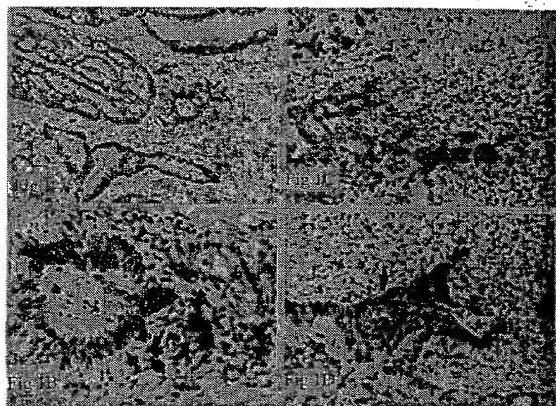


Fig. 1

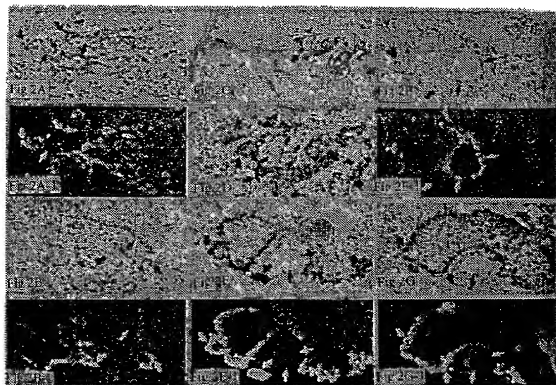


Fig. 2

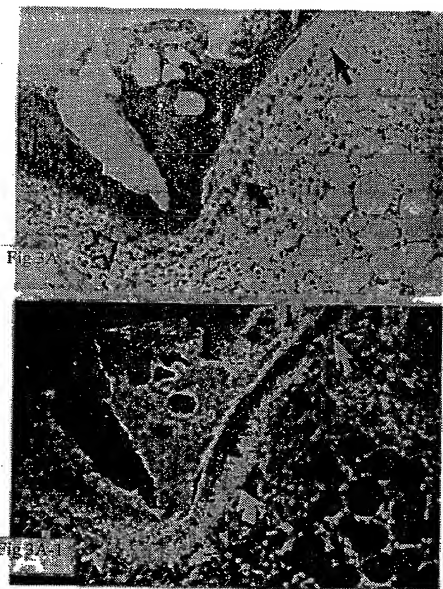


Fig. 3

FIG 4A

(SEQ ID NO.: 12 & 13)

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1 gaccacactga tcgaagggaaa aggaaggcacc agcggaagcgc agagtggagaa ccaaccaaccg
61 aggcgcggcgg cagcagacccc tgcagcggag acagagactgg agcgggccgg caccgcgcatg
1 1 M
121 cctgcgctct ggctgggctg ctcgcctctgc ttctgcctcc tctgcgccgc agcccgggcgc
2 P A L W L G C C L C F S L L L P A A R A
181 acctccaggga gggaagctctg tgattgcaat gggaagtcca ggcagtgtat ctttgatcgg
22 T S R R E V C D C N G K S R Q C I F D R
      1-0 Domain V
241 gaacttcaca gacaactcgg taatggattc cgtgcctcca actgcaatga caacactgat
42 E L H R Q T G N S F R C L N C N D N T D
301 ggcatctcact gcggaagctg caagaatggc tttaaccggc acagagaaag ggaccgcgtgt
62 G I H C E K C K N G F Y R H R E R D R C
361 ttgccttgca attgtaactc caaagttctt cttagtctct gatgtgacaa cctctggacgg
82 L P C N C N S K G S L S A R C D N S G R
421 tgcagctgta aaccagggtg gacaggagcc agatgcgacc gatgtctgcc aggcctccac
102 C S C K P G V T G A R C D R C L P G F H
481 atgtcacagg atgcgggggtg caccacaagac cagagactgc tagactccaa gtgtgactgt
122 M L T D A G C T Q D Q R L L D S K C D C
541 gaccacagctg gcatcgcagg gccctgtgac gcggggcgtg gtgtctgcaa gccagctgtt
142 D P A G I A G P C D A G R C V C K P A V
601 actggagaac gctgtgtaga gtgtcgtatca ggttactata atctgtgatgg ggggaaccct
162 T G E R C D R C R S G Y Y N L D G G N P
661 gagggctgta cccagtggtt cttgctatgg cattcagcca gctgcgcgag cctctgcagaa
182 E G C T Q C F C Y G H S A S C R S A E
      1-0 Domain IV
721 tacagtgtcc ataagatcac ctctaccttt catcaagatg ttgagggctg gaaggctgtc
202 Y S V H K I T S T F H Q D V D G W K A V
781 caacgaaatg ggtctcctgc aaagctccaa tggtcacagc gccatcaaga tgcgttttagc
222 Q R N G S P A K L Q W S Q R H Q D V F S
841 tcagcccaac gactagatcc tgtctatttt gtggctcctg ccaaatttct tgggaatcaa
242 S A Q R L D P V Y F V A P A K F L G N Q
901 caggtgagct atgggcaag cctgtccttt gactacgctg tggacagagg aggcagacag
262 Q V S Y G Q S L S F D Y R V D R G G R H
961 ccatctgccc atgatgtgat cctggaaggt gctggtctac ggatcacagc tcccttggat
282 F S A H D V I L E G A S L R I T A P L H
1021 ccacttgcca agacactgct ttgtgggctc tcaagaactt acacattcag gttcaatgag
302 P L G K T L P C G L T K T Y T F R L N E
1081 catccaagca ataattggag ccccacgctg agttactttg agtatcgaa gttactgcgg
322 H P S N N W S P Q L S Y F E Y R R L L R
1141 aatctcacag cctcccgcat ccgagctaca tatggagaat acagtactgg gtacattgac
342 N L T A L R I R A T Y G E Y S T G Y I D
1201 aatgtgaccc tgatttcagc ccgcctctgc tctggagccc cagcacccct ggttgaacag
362 N V T L I S A R P V S G A P A P W V E Q
1261 tgtatatgtc ctgttgggta caaggggcaa tctgcagag attgtgcttc tggctacag
382 C I C P V G Y K G Q F C Q D C A S G Y K
      1-0 Domain III
1321 agagattcag cgaactcggg gccttttggc acctgtattc cttgtaactg tcaaggggga
402 R D S A R L G P F G T T C I P C N C Q G G
1381 ggggctctgt atccagacac aggaagattgt tattcagggg atgagaatcc tgcattgag
422 G A C D P D T G D C Y S G D E N P D I E
1441 tgtctgact gcccaattgg ttctataaac atgcgcgacg acccccgcag ctgcaagcca
442 C A D C P I G F Y N D P H D P R S C K P
1501 tgtccctgtc ataaccgggtt cagctgctca gtgattccgg agacggagga ggtgtgtgtc
462 C P C H N G F S C S V I P E T E E V V C

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FIG 4B

(SEQ ID NO.:12 & 13)

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1561 aataaactgcc ctccccgggt caccggtgcc cgctgtgagc tctgtgtcta tggctacttt
482 N N C P P G V T G A R C E L C A D G Y F
1621 gggggccctt ttggtgaaca tggcccagtg aggccttgtc agccctgtca atgcacagc
502 G D P F G E H G P V R P C Q P C Q C N S
1681 aatgtggacc ccagtgcctc tgggaattgt gaccggctga caggcaggtg tttgaagtgt
522 N V D P S A S G N C D R L T G R C L K C
1741 atccacaaca cagccggcat ctactgcgac cagtgcgaag caggctactt cggggaccaca
542 I H N T A G I Y C D Q C K A S Y F G D P
1801 ttggctccca acccagcaga caagtgtcga gcttgcgaact gtaacccact gggtcagag
562 L A P N P A D K C R A C N C N P M G S E
1861 cctgtaggat gtccaaatga tggcacctgt gtttgcaagc caggatttgg tggcccaac
582 P V G C R S D G T C V C K P G F G G P N
1921 tgtgagcatg gacgattcag ctgtccagct tgctataatc aagtgaagat tcatgtgcat
602 C E H G A F S C P A C Y N Q V K I Q M D

      I=O Domain I/II

1981 cagtttatgc agcagcttca gagaatggag gccctgattt caaaggctca ggggtggtgat
622 Q F M Q Q L Q R M E A L I S K A Q G G D
2041 gacagatcac ctgatacaga gctggaaaggc aggatgcagc aggtctgagca ggcccttcag
642 G Y V P D T E L E G R M Q Q A E Q A L Q
2101 gacattctga gagatgccca gatttcagaa ggtgctagca gatcccttgg tctccatgtg
662 D I L R D A Q I S E G A S R S L G L Q L
2161 gccaaagtga ggagccaaga gaacagctac cagagccgctc tggatgacct caatgcagct
682 A K V R S Q E N S Y Q S R L D D L K M T
2221 gtggaaagag ttcgggtctct gggaagtccag taccagaacc gagttccggga tactcacagg
702 V E R V R A L G S Q Y Q N R V R D T H R
2281 cctcatcactc agatgcagct gagccttgcca gaaagtgaag ctctccttggg aaacactaac
722 L I T C Q H Q L S L A E S E A S L G N T N
2341 attctcgctc cagaccacta cgtggggcca aatgctctta aaagtcttggc tcaaggagcc
742 I P A S D H Y V G P N G F K S L A Q P A
2401 acaagattag cagaagaacca cgttgagtca gccattcaaca tggagcaact gacaaggaaa
762 T R L A E S H V E S A S N M E Q L T R E
2461 actgaggactc attccaaaca agccctctca ctggttgcca aggccctgca tgaaggagctc
782 T E D Y S K Q A L S L V R K A L H E G V
2521 ggaagcggaa gccgttagccc ggaagggtgct gtggttgcaag ggcttgttgg aaaaattggag
802 G S G S G S P D G A V V Q G L V E K L E
2581 aaacaccaagt ccttgggccca gcagttagaca agggaggccca ctcaagcggg aattgaagca
822 K T K S L A Q Q L T R E A T O A E I E A
2641 gataggtctt atcagcacag tctccgcttc tgctccgctc ctaggaggtc
842 D R S Y Q H S L R L L D S V S P L Q G V
2701 agtgcagctc cctttcaggt ggaagaagca aagaggatca aacaaaaagc ggattctactc
862 S D Q S F Q V E E A K R I K Q K A D S L
2761 tcaagcctgg taaccaggca tatggtatg ttcaagcgta cacaaaaaga tctgggaaac
882 S S L V T R H M D E F K R T Q K N L G N
2821 tggaaagaag aagcacagca gctcttacag attggaaaaa gtgggagaga gaaatcagat
902 W K E E A A Q Q L L Q N G K S G R E K S D
2881 cagctgcttt cccgtgccaa tcttgctaaa agcagagcac gtaagacact gagtatgggc
922 Q L L S R A N L R A K S R A L Q F A L S M G
2941 aatgccactt tttatgaagt tgagagcatc cttaaaaaac ctacagaggtt tgactgtcag
942 N A T F Y E V E S I L K N L R E F D L Q
3001 gtggacaaca gaaagcaga agctgaagaa gccattgaaga gactctctca catcagcag
962 V D N R K A E A E E A M K R L S Y I S Q
3061 aaggtttcag atgccagtga caagcccgag caagcagaaa gagcccttgg gagcgtcgt
982 K V S D A S D K T Q Q A E R A L G S A C
3121 gctgagtcac agagggcaaa gaattggggcc ggggaggccc tggaaatctc cagtgtaatt
1002 A D A Q R A K N G A G E A L E I S S E I

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FIG 4C

(SEQ ID NO.: 12 & 13)

3181 gaacaggaga ttgggagtct gaacttggaa gccaatgtga cagcagatgg agccttggcc
 1022 E Q E I G S L N L E A N V T A D G A L A
 3241 atggaaaagg gactggcctc tctgaagagt gagatgaggg aagtggaaag agagctggaa
 1042 M E K G L A S L K S E M R E V E G E L E
 3301 aggaaggagc tggagtttga cagcaatatt gatgcagttac agatgggtgat tacagaagcc
 1062 R K E L E F D T N M D A V Q M V I T E A
 3361 cagaaggttg ataccagagc caagaacgct ggggttaqaa tccaagacac acctcaacaca
 1082 Q K V D T R A K N A G V T I Q D T L N T
 3421 ttgacggccc tcttcgactct gatggaccag cctctcagtg tagatgaaga gggcgctggtc
 1102 L D G L L H L M D Q P L S V D E E G G L V
 3481 ttactggagc agaagctttc ccgagccaag acccagatca acagccaact gcggcccatg
 1122 L L E Q K L S R A K T Q I N S Q L R P M
 3541 atgtcagagc tggaaagagc ggcacgtcag cagagggggcc acctccattt gctggagaca
 1142 M S E L E E R A R Q Q R G H L H L L E T
 3601 agcatagatg ggattctggc tgatgtgaag aacttggaga acattaggga caacctggcc
 1162 S I D G I L A D V K N L E N I R D N L F
 3661 ccaggctgct acaataccca ggctctttgag caacagtga gctgccata atatttctca
 1182 F G C Y N T Q A L E Q Q

 3721 actgaggttc ttgggataca gatctcaggg ctggggagcc atgtcatgtg agtgggtggg
 3781 atggggacat ttgaacatgt ttaattggta tgctcaggtc aactgacctg accccattcc
 3841 tgatcccatg gccaggtggt tgtcttattg caccatactc ctgtcttccat gatgctgggc
 3901 atgaggcaga tagggacttg tgtgagaatg atcaaggatc tggaccccaa agatagagtg
 3961 gatggaaaga caaactgcac aggcagatgt ttgctccta atagtgttaa gtggagtctc
 4021 ggaattttga caagtgcgtg tgggatattg tcaacttatt ctttgagttaa tgtgactaaa
 4081 ggaataaact ttgactttgc ccaggcatga aattcttctt atgtlcagaa cagagtgcga
 4141 cccagtcaca ctgtggccag taataactta ttgctccta ttgtctcttg caagcttctt
 4201 gctgatcaga gtctctccta cttaacaacc aggtgtgtaa catgtctctc attttcaagc
 4261 tggaaagaat gagcagtggt ggaagtggga cctgtaagc agggccattg agagctatgg
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 4561 aaaaatgatg cgcattcaatg tattttatct tattttctca atctctctct tctttctctc
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 4681 catcttcca tccattcttc catccattac ctccatccat ccttccaca tatattatt
 4741 gagtaacctac tgtgtgccag gggctgggtg gacagtggtg acatagtctc tgccctcata
 4801 gagtgtatg tctagttagg aagacaagca ttttataaaa ataaatttaa acttacaaac
 4861 ttgtgtttgc acaagtgtg tttattgcaa taacgcgttg gtttgaagcc tctttgtcca
 4921 acagaacata tgttgcaaga ccttcccatg ggcacttgat ttggcaagac tgacagagct
 4981 ctgggtttgt cactttctt tgcatttccag cgtcactctg tgcttctcac aactgattgc
 5041 aacagactgt tgatttatga taacacagct gggaattgct ggaggaacca gggcacttcc
 5101 cactctgctt gggaaagacta tgggtctgcc ttgctctctg atttctcttg attttctgta
 5161 aagtgttttt aaataaagaa caattgttag atgccaaaaa //

FIG 4D

(SEQ ID NO.:14 & 15)

3421 ttgacggcc tctgcctct gatgggtatg tgaaccacac acccacaacc ttccagctcc
1102 L D G L L H L M G M *

3481 atgtctcagg gctttgctec agaacactca ctatacctag cccacagcaa ggggagtcct
3541 agctttcctt aaggatatca gtaaatgtgc ttgttttcca ggccacagata actttcggca
3601 ggttcacctt catttactgg accctgtttt accgttgcta agatgggtca ctgaacacct
3661 attgcacttg ggggtaaagg tctgtgggcc aaagaacagg tgtatataag caacttcaca
3721 gaacacgaga cagcttggga atcctgctaa agagtctggc ctggaccctg agaagccagt
3781 ggacagtttt aagcagagga ataacatcac cactgtatat ttcagaaaag tcactagggc
3841 agccagtttg aggaagctt gaagaggggg tttagagaaa ggcaggttga gactacttaa
3901 gatattgttg aaataattga agagagaaaat gacaggagcc tgctctaagg cagtataatg
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4021 gccgggtgtg gtggctcagc cctgtaattc tagcactttg ggagactgaa gcgggtggat
4081 cacccgaggt caggagttga agaccagcct ggccaacatg gtgaaccctt gtctctacta
4141 aaagtaaaaa aatagctgg agtgggtgg tggcgccctg taattccagc tactcaggag
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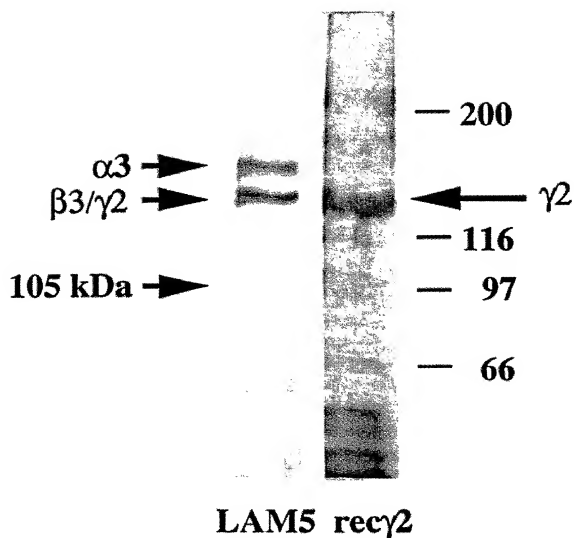


Fig. 5

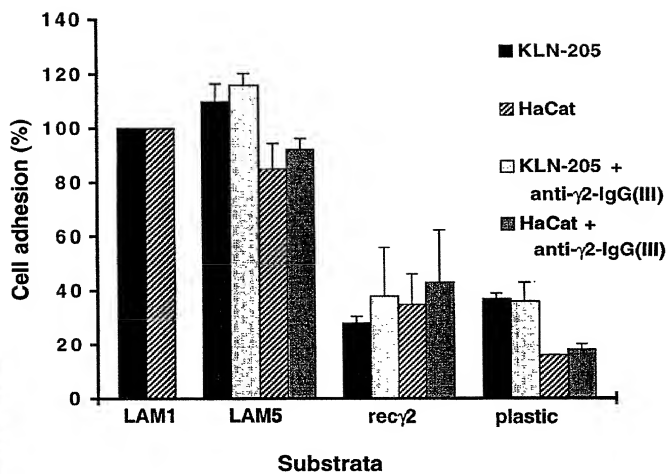


Fig. 6

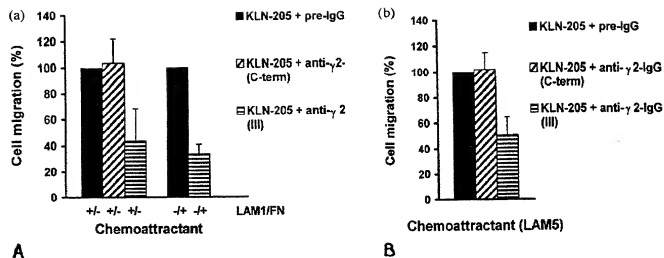


Fig 7

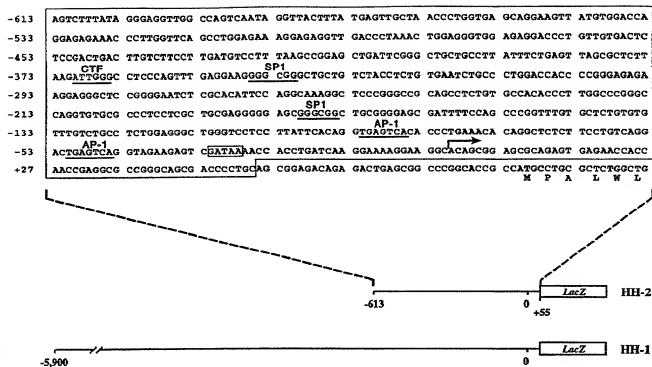


Fig. 8

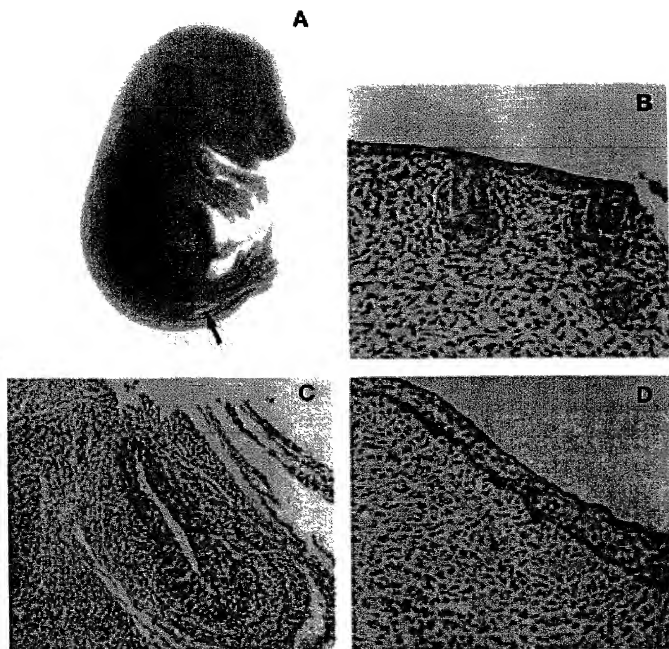


Fig. 9

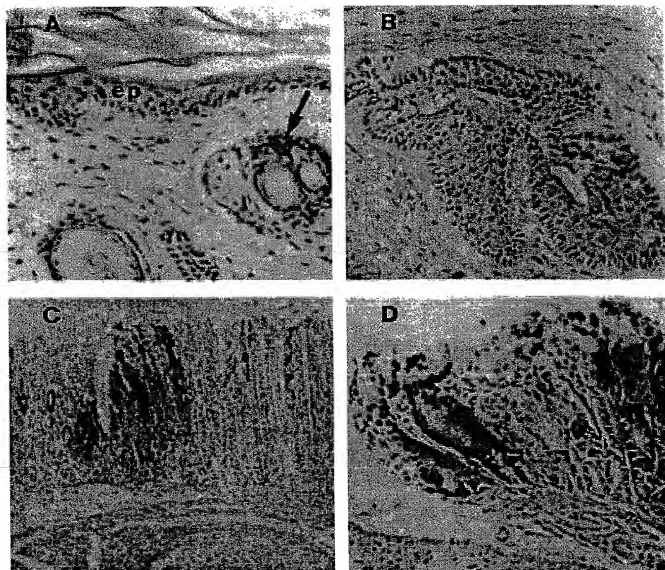


Fig. 10

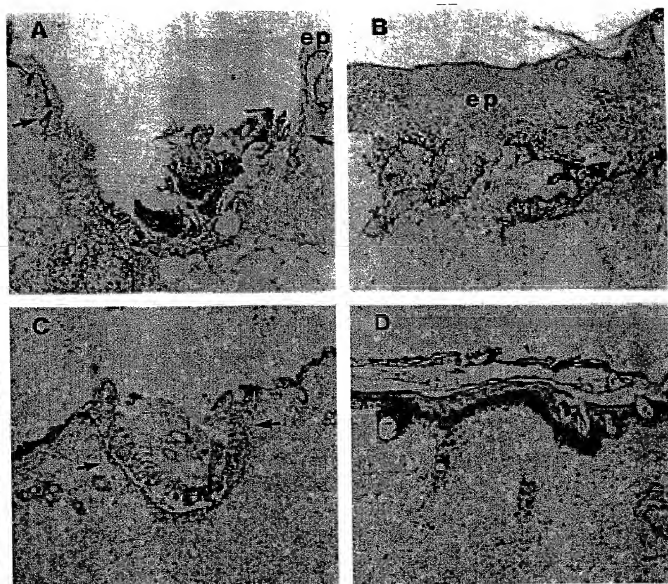


Fig. 11